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GenCore version 5.1.3
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OM protein - protein search, using sw model

January 16, 2003, 16 40-32 ; Search Lime 51 Secondo (Without alignments) 56,562 Million cell updates/see Pun on:

US-09-856-070-17 Pertect score:

1 EREKEÇMMPFKFFI. 14 Sequence:

RI CSTIMES Scoring table

Gapop 10.0 , Gapoxt 0.5

671580 Total number of hits satisfying chosen parameters:

671580 seqs, 206047115 residues

Searched.

Minimum DB seq length։ Օ Maximum DB seq length։ շըորդոր

Post-processing, Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SPTREMBL_21:* Database :

sp_archea:* sp_bacteria:* sp_tungi:*

sp_invertebrate:*
sp_mammal:* sp_organelle:* sp_human:* sp_mhc:*

sp_vertebrate:* sp_rodent:* sp_phage:* sp_plant:* sp_virus:*

sp_unclassified:*

sp_rvirus.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution. sp_bacteriap:*
sp_archeap:*

SUMMARIES

	Description	09uja2 homo sapien	Oguiz8 homo sapies	CIIIC		09uk20 homo sapien	096548 home, same	OSVBK3 rattus norv	O9vaw6 dallus dall	O9dcil mus musculu	Q18181 caenorhabdi	035763 rattus norv	O9pt71 xenopus lae	09qw27 mus sp. rad	O9pu45 dallus dall	09p219 home sapies	2950p3 drosophila
	ID	09UUZ2	Q911178	Q9H1Z7	921160	Q90K20	ชมมิงธิบั	OBVHK 3	SMS ABO	Q9DC11	018181	035763	091471	Q9QW27	091/045	Q9P219	กัจสกฤจ
	DB	-	4	4	-7	4	**	=	- 3	H	S	Π	13	Ξ	13	4	Ľ
	Query Match Length DB	152	156	158	153	161	# H =	455	585	586	556	577	580	583	583	1326	56.5
æ	Query Match	100.0	100 u	100 0	٠ ٥٠٠:	100.0	0001	95.7	95.7	95.7	73.9	73.9	73.9	73.9	73.9	73.9	68.1
	Score	99	69	69	C.	69	د ه	99	99	99	51	51	51	51	51	51	47
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qg .

RESULT 2 Q9UJZ8

Q9plz2 homo sapien Q96ju3 homo sapien	homor	Q9vyu0 drosophila		Q96pp8 homo sapien	Боло	Cadass mas masculu	OBr443 rattus norv	097116 drosophila	Q8+197 dictyosteli	Cerex mus musculu	Q9sasS dictyostell		Q8vie5 mus musculu	C9yw69 melanophus	O9yStO homo sapien	Q8wxq6 homo sapien	296p79 homo sapien	Q9usy8 schizosacch	09h6s6 homo sapien	On4466 arabidopsis	4	\$25777 plasmodium	Q9c7v7 arabidopsis	Q96pp9 homo saplen		O9p115 homo sapien	ношо
O9P1Z2 Q96JU3	060H6Ö	02770	025149	Q96PP8	Q9H253		1 Q8R443	968360	Q8T1G7	1 QAVRX2	SSONGO	1 <u>0</u> 912E6		E CHYWCG	Q9Y5TO	98 mx 06	Q96P79	8XSD6Ö	C		2 Q91FT4	025777	0 <u>0</u> 9C7V7	Q96PP9	O910W3	Q9P115	Q96T88
645 4 690 4				586 4	678 4	631 - 11	691 11		868	1241 11	1287 5	2555 11	_	365	893 4	2813 4	2817 4	166 3	~	_	_	- CO	7		730 2	793 4	793 4
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ALIGNMENTS

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Pybryczie, McLarca: Chordata; Craniata; Vertebrata, Euteleostomi;
Mammalia: Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCHI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%, Scutc 63, BH 4, Length 152:
100.0%; Pred. No. 0.0064;
tive 0, Mismatches 0, Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chen Z.C., Fadiel A., Naftolin F.;
"Mutation of cytovillin gene in glioblastoma.";
Submitted (SRP-1999) to the EMBL/GenHank/DDHJ databases.
EMBL. AF190089; AAF093158.1:
InterPro: IPR000299; Band 4.1.
InterPro: IPP000798; EZ/rad/moesin.
Pfam: PF00769; FRM: 1.
PROSIIE; PS50057; BAND_41_3; 1.
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151 AA, 18410 MW; B7FCEB175LUB9231 CRC64;
                                                                                                                              Q90JZ2;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2002 (TrEMBLrel. 20, Last annotation update)
Cylovillin 2 (Fragment).
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                                                                                       PRELIMINARY;
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Bost Local Similarity
Fine 14, Conserva'
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RESULT 1
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Best Local Similarity
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                                                          Ezrin (Fragment)
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NON_TER
SEQUENCE
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                                                                                    Homo sapieńs (Human).
Bukaryota Metanoa, Chordata; Craniata, Vertebrata; Buteleostomi,
Manmalia; Eulheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID-9606;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                          100,0%; Score 69; DB 4; Length 156;
100,0%; Pred, No. 0,0066;
Live 0; Mismatches 0; Indels
                                                                                                                                                                                  Chen Z.C., Fadiel A., Naftolin F.; "Mutation of exrin gene in cancer."; Submitted (SEP-1999) to the EMBL/GenBank/DDBJ databases. EMBL, PAF08154-1; InterPro: IPR000299; Hand_4...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chen Z.C., Fadiel A., Naftolin F.,
"Ezrin gene mutation in ovarian cancer.";
Submitted (SEP-1999) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                               156 AA; 19042 MW; 159F5AA684A2C3A6 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE 158 AA; 19086 MW; 86E92E1BC6F2957E CRC64;
                                        01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
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01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
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 PRT; 156 AA.
                            01-MAY-2000 (TrEMBLrel, 13, Created)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; AF188897; AAF03155.1; -.
InterPro; IPR000299; Band_4.1.
InterPro; IPR000798; Ez/rad/moosin.
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KT:
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NON_TER 1 1 1 1 1 NON_TER 156 156
                                                                                                                                                                                                                                                                                                                                                                        Hest Local Similarity 100.
Matches 14; Conservative
   PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                  137 EREKEÇMMREKEEL, 150
                                                                                                                                                                                                                                                                                                                                                                                                                   1 EREKEOMMREKEEL 14
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                                                                        Ezrin (Fragment).
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                                                                                                                                                                        TISSUE-BRAIN;
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NON_TER
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090JZ8;
090JZ8;
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159 AA.

PKT;

PRELLIMINARY;

1D Q9UJZ6

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                                                                                                                        Homo Sapiens (Human).
Homo Sapiens (Human).
Eukaryota: Mctazoa, Chordata, Craniata, Vertebrata, Enteleostomi:
Mammalia, Eutheria, Primates, Catarrhini: Hominidae; Homo.
NCBLTaxID-9606;
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Eukaryota; Metazoa, Chordata; Cianiata; Veitebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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"Mutation analysis of ezrin gene in cancer cells.";
"Mutation analysis of ezrin gene in cancer cells.";
Submitted (SEP-1999) to the EMBL/GenHank/DDHJ databases.
EMBL. AF18752: AAD56713.1;
InterPro: IPP000299; Band_4.1.
InterPro: IPP000789; BZ/rad/moesin.
Ptam: PF00753: Band_41; 1.
                                                                                                                                                                                                                                                                               Fadici A., Chen Z.C., Naftolin F.;
Fadici A., Chen Z.C., Naftolin F.;
Mutation of ezrin gene in cancer.";
Submitted (SEP-1999) to the EMBL/Genhank/DDHJ databases.
EMBL, AF189213: AAF03156.1;
EINGEFFC; IPR002299; Band 4.1.
InterPro: IPR00798; Ez/rad/moesin.
Pfam; PF00769; ERM; 1.
PROSITE: PS50057; HAND_41_3; 1.
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159 AA: 19234 MW: 70398388B7BA70FA CRC64;
090JZ6;
01 MAY-2000 (TYEMBLRE). 13, Created)
01-MAY-2000 (TYEMBLRE). 13, Last sequence update)
01-MAR-2002 (TYEMBLRE). 29, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-MAY-2000 (TrEMBLICEL. 13, Created)
01-MAY-2000 (TrEMBLICEL. 13, Last sequence update)
01-MAR-2002 (TrEMBLICEL. 20, Last annotation update)
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nes 14; Conservative
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Fri Jan 17 09:18:28 2003

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Mammalia: Butheria; Rodentia; Sciuroquathi; Muridae; Murinae; Pattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gales
                                                                                                                                                      Homo sapiens (Haman).
Eukaryota: Metazoa: Chordata: Craniata: Vertebrata: Eufeleostomi:
Mammalia: Eutheria: Primates: Catarrhini: Hominidae: Homo.
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100.0%; Pred No 0.021;
Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            95.7%; Score 66; DB 11; Length 455;
92.9%; Pred No 0.047;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (SEP-2001) to the EMMI/NeuBank/DDBJ databases.
EMBL: BC013903: AAH13903 1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gunn-Moore F T , Tait S , Bropby P T ;
Submitted (NOV-2001) For the EMEL/Neukruk/Roks databases
EMBL: AF450298; AAL47844.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROSITE; PSONGEG: RAND_41_1; UNKNOWN_1.
PROSITE; PSONGFI; RAND_41_2; UNKNOWN_1.
PROSITE; PSONGFI; BAND_41_3: 1.
SEQUENCE 586 AA; 69412 MW; FIB592CF49A7CC46 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   455 AA; 54174 MW; 1FC9A95F4C7D5893 CRC64;
01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-MAR 2002 (TREMBLREL. 20, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Last sequence update)
Last annotation update)
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PROSTTE; PS00661; BAND_41_2; UNKNOWN_1.
PROSTTE; PS50057; BAND_41_3; 1.
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01-MAR-2002 (TrEMBLrel, 20, Last Sequ
01-JUN-2002 (TrEMBLrel, 21, Last anno
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR060299, Band 4.1.
InterPro; IPR000799, Ez/rad/moesin.
Pfam: PF00373; Band 41, 1.
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interPro, iPR060798 Frytad/mewsin
Pfam; PF00373; Band_41; 1.
                                                                                                                      Similar to villin 2 (ezrin).
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Matches 14; Conservative
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SMART; SM00295; B41; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRELIMINARY;
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                                                                                                                                                                                                                                                                    NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                  TISSUE-COLON;
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RESULT 8

O9YGW6

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SEQUENCE FROM TISSUE-KIDNEY.

SEQUENCE FROM TISSUE-KIDNEY.

KEDLINE-21085660; PubMcd-11217851;

REDLINE-21085660; PubMcd-11217851;

RAMANI U., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Y.,

RAMANI T., Hara A., Fukunishi Y., Kondo H., Adachi J., Fukuda S.,

Alzawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,

RAGOTA K., Matsuda H.A., Ashburner M., Haraloy S., Casavant T.,

RAGOTA K., Matsuda H.A., Ashburner M., Haraloy S., Casavant T.,

RACOTA K., Matsuda H.A., Ashburner M., Bataloy S., Casavant T.,

RACOTA K., Matsuda H.S., Matsuda M., Wagner L., Washio T.,

RACOTA K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,

Blake J., Boffelli D., Bolunga N., Carninci P., de Bonaldo M.F.,

RACOMSTEIN M.J. HILL C., Hercher C., Fujira M., Gariboldi M.,

RACOMSTEIN M.J., Hoffmann M., Hune D.A., Loce N.H.,

RACOMSTEIN M.J., Hoffmann M., Hune D.A., Ramiya M., Loce N.H.,

RACOMSTEIN M.J., Makhima J., Mazzarelli J., Momberts P.,

RA Lyons P., Marchionni L., Makhima J., Mazzarelli J., Momberts P.,

RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.F.,
                                                                                                                                                                                                                                                                                               Takahashi M., Yamagata M., Noda M.;
"Specific expression of eziin, a cytoskeletal-membrane linker protein, in a subset of chick retinolectal and sensory projections.";
Eur. J. Neurosci. 11:545-558(1999).
EMBL. AND1979, BAAA7457.1, "Interpro: IPR000299; Hand.4.1.
                                                                                                                                         Eukaryo's: Metasos, Chordata, Craciata, Vertebrata; Euteleostomi,
Archosauria, Aves, Neograthao, Galliformos, Phasianidao: Phasianinao,
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Elkarysta: Mctarsa; Cherdata; Craniata; Vertebrata; Euteleostomi:
Mammalia; Eutheria; Rodentia; Seiurognathi; Muridae; Murinae; Mus.
                                                                                                                                       Chordata; Cramiata, Vertebrata; Euteleostomi;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-JUN-2001 (TrEMBLIEL 17, Created)
01-JUN-2001 (TREMBLIEL 17, Last sequence update)
01-JUN-2002 (TREMBLIEL 21, Last annotation update)
01-JUN-2002 (TREMBLIEL 11, Last annotation update)
Adult male kidney CDNA, EIKEN 1011-length enriched library,
close:0619037B12, 1011 insert sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95.7%; Score 66; DB 13; Length 585; 92.9%; Pred. No. 0.058;
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                                                                               01-MAR-2002 (Irambire). 20, mast annetation update)
                                     01-MAY-1999 (TriMBirel, 10, Created)
01-MAY-1999 (TriMBirel, 10, Last sequence update)
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  585 AA.
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STRAIN-WHITE LEGHOPN; TISSUE-BRAIN;
MEDLINE 99171943; Pubmed-10051754;
PRT;
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Ptam; PF00769; ERM; 1.
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  PRELIMINARY;
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SMAKT; SM00295; B41; 1.
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                                                                                                                    Gallus qallus (Chicken).
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    O9YGW6
                                                                                                                                                                                   Gallus
                                                                                                    Ezrin.
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Suzuki H., Toyoroka K., Wang K.H., Weitz C., Whittaker C., Wilming L., Myushaw-Boris A., Yoshida K., Haseqawa Y., Kawaji H., Kohtsuki S., Hayashizaki Y., Hayashizaki Y., "Functional annotation of a full length mouse chwa collection.",
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                                                                                                                                                                                                                                                                                                               95.7%; Score 66; DB 11; Length 586; 92.9%; Prod Ro 0.058; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             73.9%; Score 51; DB 5; Length 556; 83.3%; Pred. No. 8.7; tive 1, Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "The sequence of C. elegans cosmid C25H3.";
Submitted (JUN 1995) to the EMSL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Direct Submission.";
Submitted (SEP-2001) to the EMBL/GenBank/DDBJ databases.
EMBL: U29636: AAK31458 1; -
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Hypothetical protein.
SKOUENCE 556 AA; 61825 MW; 7D33611464040F75 CRC64;
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                                                                                                                      InterPro; IPRG00299; Band_4.1.
InterPro; IPRG00788; P2/Fad/Monsin
Pfam; PPG0378; Hand_41; 1.
Ptam; PF00769; ERM; 1.
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MFDLINE-99069613; PubMed-9851916;
                                                                                                                                                                                                                           PROSITE: PS00660; BAND_41_1; 1. PHOSITE: PS00661; BAND_41_2; 1. PHOSITE: PS50057; BAND_41_3; 1. SEQUENCE: SBE AA: DA44 MW:
                                                                                     EMBL; AK002766; BAB22341.1; -.
MGD; MGI:98931; Vil2.
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Best Local Similarity 92 99
"*Febres 13; Conservative
                                                                       Nature 409:685-690(2001).
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Johnson D., Wilson R.;
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Best Local Similarity
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                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia: Eutheria; Eodentia; Sciurognathi, Muridae: Murinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota, Metažča, Chordata, Craniata, Vertebrata, Euteleostomi;
Amphibia, Batrachia, Anura, Meschatrachia, Fipoidea: Pipidae;
Xenopodinae; Xenopus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Indels
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Submitted (JUN-1995) to the EMBL/GenBank/DDBJ databases.
EMBL; U29763; AARLILLS.;
InterPro; IPPOND299; Band_4.1.
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PROSITE, PROUGES, B41, 1
PROSITE, PSOUGE1; BAND_41_2; 1.
PROSITE; PSSUUS7; BAND_41_2; 1.
PROSITE; PSSUUS7; BAND_41_3; 1.
SEQUENCE 580 AA; b8134 MW; 100bA7E9E5887074 CRC64;
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                                                                                                                             01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01 MAR 2002 (TrEMBLrel. 20, Last annotation update)
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20, Last asnotation update)
                                                577 AA
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01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last seq
01-MAP-2002 (TrEMBLrel 20, Last ann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Xenopus laevis (African clawed frog).
                                                                                                   (reated)
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                                                     PPT;
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PROSITE: PS00661: BAND 41_2: 1.
PROSITE: PS50057: BAND 41_3: 1.
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Pfam; PF00769; ERM; 1.
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                                                   PRF1 IMINARY;
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                                                                                                                                                                                                  Rattus norvegicus (Rat).
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RESULT 11
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
      0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "The ezrin-like family of Lyrosine kinase substrates: receptor-specific pattern of Lyrosine phosphorylation and relationship to malignant transformation."; Oncogene 8 1335-1345(1349).
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Fazioli F., Wong W.T., Hilrich S.J., Sakaguchi K., Appella F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              73.9%, Score 51, DR 11, Langth 583, 71.4%; Pred. No. 9.1;
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Li W., Crouch D.H.;
"Clounting and expression profile of chicken radixin.";
Blochim Piophys, Acta 1491:327-332(2000).
EMBL; AJ249838; CABS9977.1; -.
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Last annotation update)
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01-MAY-2000 (TrEMBLrel. 13, Last. sequence update)
01-MAR-2002 (TrEMBLrel. 26, Last annotation update)
MADIXIN-ESP10 product.
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01-MAR-2002 (TYEMBLEEL: 20, Last anno
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InterPro; 1PR00798; E2Z/ad/MocesIn.
Plam: PF0073; Band_41; 1.
Plam: PF00769; ERM: 1.
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InterPro, IPR000798: Ez/rad/mnosin
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Pfam: PF00769: ERM: 1.
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SMART; SM00295; B41; 1.
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                                                          1 BREKEÇMMREKEEL, 14
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EMBL. ABD49:42. BAN9:933.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens (Human).
Eukarycta, Metazoa, Chordata, Craniata; Vertebrata, Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID-9606;
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                                                                                                                  73.9%; Score 51; DB 13; Length 583; 71.4%; Pred. No. 9.1;
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                                                                                                                                                      1; Indels
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                            PROSITE; PS00660; BAND_41_1; 1.
PROSITE; PS00661; BAND_41_2; 1.
PROSITE; PS050057; HAND,41_3; 1.
SEQUENCE PS3 AA: 68555 MW; BE25634F4798CBE0 CRC64;
                                                                                                                                                                                                                                                                                                                                        01-0CT-2000 (FFBMEREL, 15, Created)
01-0CT-2000 (FFBMEREL, 15, Last sequence update)
01-0CT-2000 (FFBMEREL, 15, Last annotation update)
                                                                                                                                                                                                                                                                                                          PRT; 1326 AA
                                                                                                                                                      3; Mismatches
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nes 10, Conservative
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PRINTS; PR00935; BAND41.
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                 SMART; SM00295; B41;
                                                                                                                                      Local Similarity
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